· To: Ms. Minu yi'm

AS



OIPE

RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/813,153

DATE: 07/11/2002 TIME: 13:06:27

Input Set : N:\Crf3\RULE60\09813153.raw
Output Set: N:\CRF3\07112002\1813153.raw

1 <110> APPLICANT: Rosen et al.
2 <120> TITLE OF INVENTION: 67 Human secreted proteins

3 <130> FILE REFERENCE: PZ023 4 <140> CURRENT APPLICATION NUMBER: 09/813,153

5 <141> CURRENT FILING DATE: 2001-03-21

6 <150> PRIOR APPLICATION NUMBER: US/09/363,044

7 <151> PRIOR FILING DATE: 1999-07-29

8 <150> PRIOR APPLICATION NUMBER: 60/073,160

9 <151> PRIOR FILING DATE: 1998-01-30

10 <150> PRIOR APPLICATION NUMBER: 60/073,159

11 <151> PRIOR FILING DATE: 1998-01-30

12 <150> PRIOR APPLICATION NUMBER: 60/073,165

13 <151> PRIOR FILING DATE: 1998-01-30

14 <150> PRIOR APPLICATION NUMBER: 60/073,164

15 <151> PRIOR FILING DATE: 1998-01-30

16 <150> PRIOR APPLICATION NUMBER: 60/073,167

17 <151> PRIOR FILING DATE: 1998-01-30

18 <150> PRIOR APPLICATION NUMBER: 60/073,162

19 <151> PRIOR FILING DATE: 1998-01-30

20 <150> PRIOR APPLICATION NUMBER: 60/073,161

21 <151> PRIOR FILING DATE: 1998-01-30

22 <150> PRIOR APPLICATION NUMBER: 60/073,170

23 <151> PRIOR FILING DATE: 1998-01-30

24 <160> NUMBER OF SEQ ID NOS: 298

25 <170> SOFTWARE: PatentIn Ver. 2.0

27 <210> SEQ ID NO: 1

28 <211> LENGTH: 733

29 <212> TYPE: DNA

30 <213> ORGANISM: Homo sapiens

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35		tcaagttcaa	ctggtacgtg	gacggcgtgg	aggtgcataa	tgccaagaca	aagccgcggg	240
36		aggagcagta	caacagcacg	taccgtgtgg	tcagcgtcct	caccgtcctg	caccaggact	300
37		ggctgaatgg	caaggagtac	aagtgcaagg	tctccaacaa	agccctccca	acccccatcg	360
38		agaaaaccat	ctccaaagcc	aaagggcagc	cccgagaacc	acaggtgtac	accctgcccc	420
39		catcccggga	tgagctgacc	aagaaccagg	tcagcctgac	ctgcctggtc	aaaggcttct	480
40		atccaagcga	catcgccgtg	gagtgggaga	gcaatgggca	gccggagaac	aactacaaga	540
41		ccacgcctcc	cgtgctggac	tccgacggct	ccttcttcct	ctacagcaag	ctcaccgtgg	600
42		acaagagcag	gtggcagcag	gggaacgtct	tctcatgctc	cgtgatgcat	gaggctctgc	660
43		acaaccacta	саспсапаап	agentetece	tatetecaaa	taaatgagtg	cgacggccgc	720

ENTERED

733

44

Input Set : N:\Crf3\RULE60\09813153.raw
Output Set: N:\CRF3\07112002\1813153.raw

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                                                                                        120
    80
              gcccctaact ccgcccaftt ccgcccattc tccgccccat ggctgactaa ttttttttat
                                                                                        180
              ttatgcagag gccgaggccg cctcggcctc tgagctattc cagaagtagt gaggaggctt
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                                                                                        240
    82
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Output Set: N:\CRF3\07112002\I813153.raw

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120		cagttccgcc cattctccgc cccatggctg actaattttt tttatttatg cagaggccga	180					
121		ggccgcctcg gcctctgagc tattccagaa gtagtgagga ggcttttttg gaggcctagg	240					
122	.010	cttttgcaaa aagctt	256					
		SEQ ID NO: 11						
		LENGTH: 1079						
		TYPE: DNA ORGANISM: Homo sapiens						
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129	\400 /	ggcacgagcc aatttgccaa ggttctaaag gcttatgagg tcctgaagga gccaggcctt	60					
130		gtgatggagt aggtgacaca ggcctggttg tcctgtcagc agaagggaaa gcaggggctg	120					
131		ggctgagagg aggacacgga gggctctgct gaggttcctt cctgggttcc accaacaggg	180					
132		acagggagte acttgeette cagttetgtg etgggatgge gggacageae ttggettget	240					
133		tggccagctg cgtcatgagt ttgatttggt ttttttttt ttgcagctgc ttcatatgct	300					
134		ctgctccagc ccctccccaa cagctggtag cttatggttt cttcaagagg aaagtagact	360					
135		ttatgctgta catttgagct gtagagctaa gattcgctta ctggtgagct gtgaaacctt	420					
136		gttgcttttt cccagagtct gatggcagtg actgtgatca agggaatctt caccgccaca	480					
137		agtgcaggca gcaggtgtgg ttcaggtccc ccccacccc actgtgctcc tttgaagcca	540					
138		acgtgcctcc ctcgcctcca tactggaggg acgacgcagg ggagaacaga gaagtgcttg	600					
139		gccctaggat tgaggcactt gtttcctagc ccgctgggtt agggctggtg caagcgaggc	660					
140		aatgttgagg atgctttaag cactaccagc cgaatccggg aactctgtta acagttgtcc	720					
141		aaccagcaga atgaggctaa ctgtataaag catgggaccc aggatgagga taaggaaagg	780					
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143		gactgtgact ctggtgagga ggggtgagca gggaggttga ttctctgatg ttaactaagt	900					
144		ggcaaagtct caaccgtgct cagccctccc cctcccaggg aagagaaaca aagattcaaa	960					
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Input Set : N:\Crf3\RULE60\09813153.raw
Output Set: N:\CRF3\07112002\I813153.raw

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155		ctgcctcagc	accatggtgc	gccaggtccc	gacggctccg	cgccagatcc	cgcccactac	180
156		agggagcgag	tcaaggccat	gttctaccac	gcctacgaca	gctacctgga	gaatgccttt	240
157		cccttcgatg	agctgcgacc	tctcacctgt	gacgggcacg	acacctgggg	cagtttttct	300
158		ctgactctaa	ttgatgcact	ggacaccttg	ctgattttgg	ggaatgtctc	agaattccaa	360
159		agagtggttg	aagtgctcca	ggacagcgtg	gactttgata	ttgatgtgaa	cgcctctgtg	420
160				ggtaggagga				480
161		gctggggtgg	aagtagaggc	tggatggccc	tgttccgggc	ctctcctgag	aatggctgag	540
162				cccagccttt				600
163				gaacccagga				660
164				tgccaccctg				720
165				gatgcgcctc				780
166				cactggcaag				840
167				gtacttggtg				900
168				gtataacaaa				960
169				gtacaagggg				1020
170				kcagagcctc				1080
171				atggaagcag				1140
172				gaagcgagag				1200
173				tgccacgggg				1260
174				aatcagcaag				1320
175				caaccgcatg				1380
176				cccrrccaac				1440
177				ggagtgcatc				1500
178				tgccgccctg				1560
179				gagggaattc				1620
180				ggggccatgg				1680
181				ggcaagggag				1740
182				cttcacctcc				1800
183				taatttttt				1860
184				aaaaaaaaa				1920
185		aagggcggcc						1932
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		SEQUENCE: 1						
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193				atgactatgg				120
194			-	cggtgggccc				180
195				agggaattga				240
196				aaagtatatt				300
197				tggacttgaa				360
198		-		ttttttgcat				420
199			-	ggttttagag				480
200			-	ggaagagccg		-		540
201			_	gctgcctccc				600
202				aggcgctcac				660
203				ccctctgctt				720
			,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,		-555544490			0

Input Set : N:\Crf3\RULE60\09813153.raw
Output Set: N:\CRF3\07112002\1813153.raw

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205	catcagggcg tgaaatcaaa ctagatgtgg gcagg	ggagag kgttgcttac ctgcctgctg	840
206	gggcagggtt tcctgaaact gggttaattc ttta	tagaaa tgtgaacact gaatttattt	900
207	taaaaaataa taataaaaat ttaaaaaaat taaaa	aataaa aaaaaccaca gaaaacaact	960
208	ttacatgtat ataggtcttg aagtgagtga agtg	gctgct tttttttttt tttttttt	1020
209	gcttttttt gctttttgta gaagagattg agaat	tggtac tctaatcaaa aataaagttt	1080
210	tgtagtggga ccagaaatta cttacctgac atcca		1140
211	ggttgaaagt tocagacotg ctgtcgaggc cttg		1200
212	tgcaaggacg caactgtgag ctgaggtgtg agcc		1260
213	ccqctqctqc caqcctcaqa aaggcaccca ggtg		1320
214	gccccagga atcaaggata gggctaaggt tttca		1380
215	aggtgactgc ttcctcccgc ccttcacaga actga		1440
216	agggggccgg ggctcaggag ccatgacctg gtgt		1500
217	atgtgaatgg agacaggtat gagaggctgt cotog		1560
218	togggootca ogacggtgot acctaagaaa gtot		1620
210			1680
	cagtggtcag caaattggaa gaggatccga tggga		1740
220	gattatacct gtttgtggtt tagctttgta ttta		1800
221	ttgtcatcat aaaaatgaaa caaattaaaa tatti	tattyc caggcaaaaa aaaaaaaaa	
222	aaaaaaaaa aaaaaaaaa aaaaaaa		1827
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	LENGTH: 696		
	TYPE: DNA		
	ORGANISM: Homo sapiens		
	SEQUENCE: 14		
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230	tgggagaatt tacttctgtg gtttgctact gtttc		120
231	ttgttaggtg gcagggttgt ggggcagagt ggggt		180
232	agagggcaca ggaagatctg taagcaagag ggcat		240
233	ggaaggttga caggccaaag aaagatcagc teete		300
234	togtagoooc agacacgttt totcaatttg agcac	caatat ccattactat ttcccgtact	360
235	gggtttcaat taaagagagt gagagtagaa agtto	cactgg tgtttggggg ttcatttatt	420
236	tccaagcagg atgcaaatga aagggagccg tggg	cacaga gttgtcatgt gtgtttttcc	480
237	tecetettet ttecatttee ttettgeaac ettec	ceteca ettettgeca gecacecage	540
238	acacccgtgt tcccaaagca aatgttttca wgtct	ttgaaa atccagttag ggtgaggaga	600
239	gaaggaaggt gataacatca tacctactga tgcco	ccctag agatgaagct gtcctggggg	660
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	LOCATION: (736)		
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252	tgctaaggtt ctgtgctaca aattgatgtt tagat		120
253	gaattggtgc ataaggctgc ctcctgytat atgg		180
254	tgcaacaacc agcttccagt ttactacacc tacaa		240
	- · · · · · · · · · · · · · · · · · · ·		

Input Set : N:\Crf3\RULE60\09813153.raw
Output Set: N:\CRF3\07112002\I813153.raw

Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

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Seq#:15; N Pos. 736
Seg#:20; N Pos. 547,555
Seq#:21; N Pos. 396
Seq#:22; N Pos. 48,987,1037,1312
Seq#:23; N Pos. 720
Seq#:25; N Pos. 910,912,958,1038
Seq#:26; N Pos. 15,27
Seq#:27; N Pos. 2,13,61,79,101
Seq#:28; N Pos. 1063,1067
Seq#:33; N Pos. 184,208
Seq#:34; N Pos. 9,1692
Seq#:36; N Pos. 13
Seq#:40; N Pos. 38
Seq#:42; N Pos. 770
Seg#:43; N Pos. 798,927,944,974,1035,1058
Seq#:47; N Pos. 488,536,539,548
Seq#:51; N Pos. 1874
Seq#:52; N Pos. 5,136
Seq#:53; N Pos. 502
Seq#:57; N Pos. 353,528
Seq#:58; N Pos. 1480
Seq#:67; N Pos. 76
Seq#:69; N Pos. 323,337
Seq#:75; N Pos. 136
Seq#:78; N Pos. 1212
Seq#:79; N Pos. 9,410
Seq#:81; Xaa Pos. 54
Seq#:82; Xaa Pos. 326,342,444
Seq#:83; Xaa Pos. 100
Seq#:84; Xaa Pos. 48
Seq#:85; Xaa Pos. 47
Seq#:86; Xaa Pos. 276
Seq#:87; Xaa Pos. 86
Seq#:88; Xaa Pos. 95
Seq#:89; Xaa Pos. 313
Seq#:90; Xaa Pos. 80
Seq#:91; Xaa Pos. 47
Seq#:92; Xaa Pos. 106,129
Seq#:93; Xaa Pos. 71
Seq#:94; Xaa Pos. 48
Seg#:95; Xaa Pos. 57
Seg#:96; Xaa Pos. 73
Seq#:97; Xaa Pos. 60
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Input Set : N:\Crf3\RULE60\09813153.raw
Output Set: N:\CRF3\07112002\I813153.raw

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Seq#:101; Xaa Pos. 83
Seq#:102; Xaa Pos. 43
Seg#:103; Xaa Pos. 286,318
Seq#:104; Xaa Pos. 46
Seq#:105; Xaa Pos. 49
Seq#:106; Xaa Pos. 66
Seq#:107; Xaa Pos. 46
Seq#:108; Xaa Pos. 74
Seq#:109; Xaa Pos. 50
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Seq#:111; Xaa Pos. 74
Seq#:112; Xaa Pos. 43
Seq#:113; Xaa Pos. 180
Seq#:114; Xaa Pos. 47
Seq#:115; Xaa Pos. 34,81
Seq#:116; Xaa Pos. 68
Seg#:117; Xaa Pos. 43
Seq#:118; Xaa Pos. 212
Seq#:119; Xaa Pos. 44
Seq#:120; Xaa Pos. 127
Seq#:121; Xaa Pos. 57
Seq#:122; Xaa Pos. 46
Seq#:123; Xaa Pos. 47
Seq#:124; Xaa Pos. 57
Seq#:125; Xaa Pos. 273
Seq#:126; Xaa Pos. 281
Seq#:127; Xaa Pos. 83,141
Seq#:128; Xaa Pos. 188,211,295
Seq#:129; Xaa Pos. 43
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Seq#:131; Xaa Pos. 427
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Seq#:137; Xaa Pos. 63
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Seq#:141; Xaa Pos. 221
Seq#:142; Xaa Pos. 60
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Seq#:144; Xaa Pos. 59
Seq#:145; Xaa Pos. 68
Seq#:146; Xaa Pos. 177
Seq#:147; Xaa Pos. 120
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Input Set : N:\Crf3\RULE60\09813153.raw
Output Set: N:\CRF3\07112002\1813153.raw

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Seq#:158; Xaa Pos. 9,31
Seq#:159; Xaa Pos. 12
Seq#:161; Xaa Pos. 37
Seq#:162; Xaa Pos. 7
Seq#:165; Xaa Pos. 6,56,67,113,130,137,139,211,222,224,227,280
Seq#:166; Xaa Pos. 6
Seq#:167; Xaa Pos. 12,23
Seq#:168; Xaa Pos. 27,44
Seg#:169; Xaa Pos. 5,7
Seq#:170; Xaa Pos. 38
Seq#:171; Xaa Pos. 5,7,10
Seq#:172; Xaa Pos. 20
Seq#:178; Xaa Pos. 309,341
Seq#:186; Xaa Pos. 2,34
Seq#:197; Xaa Pos. 236,324
Seq#:203; Xaa Pos. 13
Seq#:205; Xaa Pos. 9
Seq#:217; Xaa Pos. 56
Seq#:219; Xaa Pos. 9
Seq#:221; Xaa Pos. 84
Seq#:222; Xaa Pos. 118,176
Seq#:255; Xaa Pos. 185
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Invalid Line Length:

The rules require that a line not exceed 72 characters in length. This includes spaces.

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Seq#:3; Line(s) 63,64
Seq#:4; Line(s) 71
Seq#:5; Line(s) 78,79,80,81,82
Seq#:6; Line(s) 89
Seq#:7; Line(s) 96
Seq#:8; Line(s) 103
Seq#:9; Line(s) 110,111
Seq#:10; Line(s) 118,119,120,121,122
Seq#:11; Line(s) 129,130,131,132,133,134,135,136,137,138,139,140,141,142
Seq#:11; Line(s) 143,144,145,146
Seq#:12; Line(s) 153,154,155,156,157,158,159,160,161,162,163,164,165,166
Seq#:12; Line(s) 167,168,169,170,171,172,173,174,175,176,177,178,179,180
Seq#:12; Line(s) 181,182,183,184,185
Seq#:13; Line(s) 192,193,194,195,196,197,198,199,200,201,202,203,204,205
Seq#:13; Line(s) 206,207,208,209,210,211,212,213,214,215,216,217,218,219
Seq#:13; Line(s) 220,221,222
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